IFW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/072,301A

DATE: 08/06/2004
TIME: 11:22:50

Input Set : A:\PTO.LM.txt
Output Set: N:\CRF4\08062004\J072301A.raw

```
 3 <110> APPLICANT: Hua, Shaobing
 4         Pauling, Michelle H.
 5         Zhu, Li
 7 <120> TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CORECEPTORS FOR HUMAN
 8         IMMUNODEFICIENCY VIRUS
10 <130> FILE REFERENCE: 25636-718
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C--> 12 **<140> CURRENT APPLICATION NUMBER: US/10/072,301A**
C--> 12 **<141> CURRENT FILING DATE: 2002-02-08**

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12 <160> NUMBER OF SEQ ID NOS: 60
14 <170> SOFTWARE: PatentIn version 3.1
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 352
18 <212> TYPE: PRT
19 <213> ORGANISM: Homo sapiens
21 <400> SEQUENCE: 1
23 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
24 1               5                   10                  15
27 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
28              20                  25                  30
31 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
32          35                  40                  45
35 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
36      50                  55                  60
39 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
40 65                  70                  75                  80
43 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
44                  85                  90                  95
47 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
48              100                 105                 110
51 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
52          115                 120                 125
55 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
56      130                 135                 140
59 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
60 145                 150                 155                 160
63 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
64              165                 170                 175
67 Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn
68              180                 185                 190
71 Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu
72          195                 200                 205
75 Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys
76      210                 215                 220
```

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```
79 Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile
80 225             230             235                     240
83 Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu
84             245             250                     255
87 Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser
88             260             265                     270
91 Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr
92         275             280                 285
95 His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe
96     290             295                 300
99 Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe
100 305             310             315                     320
103 Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser
104             325             330                     335
107 Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu
108             340             345                 350
111 <210> SEQ ID NO: 2
112 <211> LENGTH: 17
113 <212> TYPE: PRT
114 <213> ORGANISM: Homo sapiens
116 <400> SEQUENCE: 2
118 Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser Asn Arg Leu Asp
119 1             5                 10                 15
122 Gln
126 <210> SEQ ID NO: 3
127 <211> LENGTH: 32
128 <212> TYPE: PRT
129 <213> ORGANISM: Homo sapiens
131 <400> SEQUENCE: 3
133 Thr Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser His Phe
134 1             5                 10                 15
137 Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile
138             20             25                 30
141 <210> SEQ ID NO: 4
142 <211> LENGTH: 20
143 <212> TYPE: PRT
144 <213> ORGANISM: Artificial Sequence
146 <220> FEATURE:
147 <223> OTHER INFORMATION: G4S Linker
149 <400> SEQUENCE: 4
151 Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
152 1             5                 10                 15
155 Gly Gly Gly Ser
156             20
159 <210> SEQ ID NO: 5
160 <211> LENGTH: 60
161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial Sequence
164 <220> FEATURE:
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Input Set : **A:\PTO.LM.txt**
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```
165 <223> OTHER INFORMATION: DNA of G4S Linker
167 <400> SEQUENCE: 5
168 ggcggtggtg gatcaggcgg cggaggatct ggcggaggtg gcagcggtgg tggaggcagt      60
171 <210> SEQ ID NO: 6
172 <211> LENGTH: 60
173 <212> TYPE: DNA
174 <213> ORGANISM: Artificial Sequence
176 <220> FEATURE:
177 <223> OTHER INFORMATION: 5' Homologous Sequence
179 <400> SEQUENCE: 6
180 accccaccaa acccaaaaaa agagatctgt atggcttacc catacgatgt tccagattac      60
183 <210> SEQ ID NO: 7
184 <211> LENGTH: 57
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
189 <223> OTHER INFORMATION: 3' Homologous Sequence
191 <400> SEQUENCE: 7
192 gagatggtgc acgatgcaca gttgaagtga acttgcgggg tttttcagta tctacga        57
195 <210> SEQ ID NO: 8
196 <211> LENGTH: 36
197 <212> TYPE: PRT
198 <213> ORGANISM: Homo sapiens
200 <400> SEQUENCE: 8
202 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
203 1               5                   10                  15
206 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
207             20                  25                  30
210 Leu Pro Pro Leu
211         35
214 <210> SEQ ID NO: 9
215 <211> LENGTH: 32
216 <212> TYPE: PRT
217 <213> ORGANISM: Homo sapiens
219 <400> SEQUENCE: 9
221 Thr Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser His Phe
222 1               5                   10                  15
225 Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile
226             20                  25                  30
229 <210> SEQ ID NO: 10
230 <211> LENGTH: 30
231 <212> TYPE: DNA
232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Primer
237 <400> SEQUENCE: 10
238 ggagaattcg attatcaagt gtcaagtcca                                      30
241 <210> SEQ ID NO: 11
242 <211> LENGTH: 31
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Input Set : **A:\PTO.LM.txt**
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```
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial Sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: Primer
249 <400> SEQUENCE: 11
250 cgcggatcct tagagcggag gcaggaggcg g                          31
253 <210> SEQ ID NO: 12
254 <211> LENGTH: 29
255 <212> TYPE: DNA
256 <213> ORGANISM: Artificial Sequence
258 <220> FEATURE:
259 <223> OTHER INFORMATION: Primer
261 <400> SEQUENCE: 12
262 ggagaattca ccagatctca aaaagaagg                             29
265 <210> SEQ ID NO: 13
266 <211> LENGTH: 33
267 <212> TYPE: DNA
268 <213> ORGANISM: Artificial Sequence
270 <220> FEATURE:
271 <223> OTHER INFORMATION: Primer
273 <400> SEQUENCE: 13
274 cgcggatcct tatatcttta atgtctggaa att                        33
277 <210> SEQ ID NO: 14
278 <211> LENGTH: 21
279 <212> TYPE: DNA
280 <213> ORGANISM: Artificial Sequence
282 <220> FEATURE:
283 <223> OTHER INFORMATION: Primer
285 <400> SEQUENCE: 14
286 caggaattct ttggcctgaa t                                     21
289 <210> SEQ ID NO: 15
290 <211> LENGTH: 31
291 <212> TYPE: DNA
292 <213> ORGANISM: Artificial Sequence
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Primer
297 <400> SEQUENCE: 15
298 cgcggatcct cagcagtgcg tcatcccaag a                          31
301 <210> SEQ ID NO: 16
302 <211> LENGTH: 759
303 <212> TYPE: DNA
304 <213> ORGANISM: Artificial Sequence
306 <220> FEATURE:
307 <223> OTHER INFORMATION: Clone 15.186.35
309 <400> SEQUENCE: 16
310 caggttacct tgaaggagtc tggtcctacg ttggtgaaac ccacacagac cctcacgctg    60
312 acctgcacct tgtctgggtt ctcactcagc actagtggag tgagtgtggg ctggatccgt   120
314 cagcccccag gaaaggccct tgagtggctt gcaagcataa attggaatga tgataagtgc   180
316 tacagcccat ctctgaaaag caggctcacc atcaccaagg acaccccaa  aaaccaggtg   240
```

```
318 gtccttgcaa tgagcaacat ggaccctgcg gacacagcca catattcctg tgcactcgat   300
320 atgccccccc atgatagtgg cccgcaatct tttgatgctt ctgatgtctg gggcccaggg   360
322 acaatggtca ccgtctcttc aggcggtggt ggatcaggcg gcggaggatc tggcggaggt   420
324 ggcagcggtg gtggaggcag ttcctatgag ctgatgcagc taccctcagt gtccgtgtcc   480
326 ccaggacaga cagccagcat cacctgctct ggagataatt tgggggataa atatgcctgc   540
328 tggtatcaac agaagccagg ccggtcccct gtgctggtca tttatggaga taacaagcgg   600
330 ccctcaggga tccctgagcg attctctggc tccaactctg gaacacagc cactctgacc    660
332 atcagcggga cccaggctat ggatgaggct gactattact gtcaggcgtg ggacaccagc   720
334 actgctgtct cggaactgg gaccaagctc accgtccta                          759
337 <210> SEQ ID NO: 17
338 <211> LENGTH: 253
339 <212> TYPE: PRT
340 <213> ORGANISM: Artificial Sequence
342 <220> FEATURE:
343 <223> OTHER INFORMATION: Clone 15.186.35
345 <400> SEQUENCE: 17
347 Gln Val Thr Leu Lys Glu Ser Gly Pro Thr Leu Val Lys Pro Thr Gln
348 1               5                   10                  15
351 Thr Leu Thr Leu Thr Cys Thr Leu Ser Gly Phe Ser Leu Ser Thr Ser
352             20                  25                  30
355 Gly Val Ser Val Gly Trp Ile Arg Gln Pro Pro Gly Lys Ala Leu Glu
356         35                  40                  45
359 Trp Leu Ala Ser Ile Asn Trp Asn Asp Asp Lys Cys Tyr Ser Pro Ser
360     50                  55                  60
363 Leu Lys Ser Arg Leu Thr Ile Thr Lys Asp Thr Pro Lys Asn Gln Val
364 65                  70                  75                  80
367 Val Leu Ala Met Ser Asn Met Asp Pro Ala Asp Thr Ala Thr Tyr Ser
368             85                  90                  95
371 Cys Ala Leu Asp Met Pro Pro His Asp Ser Gly Pro Gln Ser Phe Asp
372             100                 105                 110
375 Ala Ser Asp Val Trp Gly Pro Gly Thr Met Val Thr Val Ser Ser Gly
376         115                 120                 125
379 Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
380     130                 135                 140
383 Gly Gly Ser Ser Tyr Glu Leu Met Gln Leu Pro Ser Val Ser Val Ser
384 145                 150                 155                 160
387 Pro Gly Gln Thr Ala Ser Ile Thr Cys Ser Gly Asp Asn Leu Gly Asp
388             165                 170                 175
391 Lys Tyr Ala Cys Trp Tyr Gln Gln Lys Pro Gly Arg Ser Pro Val Leu
392             180                 185                 190
395 Val Ile Tyr Gly Asp Asn Lys Arg Pro Ser Gly Ile Pro Glu Arg Phe
396         195                 200                 205
399 Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr
400     210                 215                 220
403 Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp Asp Thr Ser
404 225                 230                 235                 240
407 Thr Ala Val Phe Gly Thr Gly Thr Lys Leu Thr Val Leu
408             245                 250
411 <210> SEQ ID NO: 18
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/06/2004
PATENT APPLICATION: **US/10/072,301A**   TIME: 11:22:51

Input Set : **A:\PTO.LM.txt**
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:32; Xaa Pos. 4
Seq#:33; Xaa Pos. 3,4

VERIFICATION SUMMARY   DATE: 08/06/2004
PATENT APPLICATION: **US/10/072,301A**   TIME: 11:22:51

Input Set : **A:\PTO.LM.txt**
Output Set: **N:\CRF4\08062004\J072301A.raw**

```
L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:1222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
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